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d) insertion, e) site-directed mutagenesis or f) naturally occurring mutation that causes a loss of biological function in the *NAP1L2* gene or *Nap1l2* gene.

61. (NEW) The neural cell of claim 57, wherein the *NAP1L2* gene is modified, wherein the modification is selected from a) substitution, b) deletion, c) frameshift, d) insertion, or e) site-directed mutagenesis that causes a loss of biological function in the *NAP1L2* gene.

**REMARKS**

Entry of this Amendment prior to examination is respectfully requested.

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The amendment to page 6 of the specification does not add new matter because it merely corrects a typographical error. A "Napsll" gene is not the focus of the invention, and is not recited anywhere else in the specification. Instead, the term "Napsll" is a typographical error in the term "*Nap1l2*," a term which is recited numerous other times in the specification. The error in the term "Napsll" is also clear because it is used on page 6 in conjunction with the term "*NAP1L2*", which is the human homologue of the mouse *Nap1l2* gene.

New claims 54-61 and amended claims 1-3, 5-7, 9, 15, 17, 19, 20, 22, 24, 26, 27, 28, 33-37, 39, 41, 45 and 46 do not add new matter to the specification, but were derived from the original claims and were changed to conform to United States patent practice.

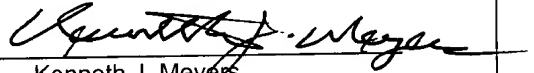
If there is any fee due in connection with the filing of this Preliminary Amendment, please charge the fee to our Deposit Account No. 06-0916.

Respectfully submitted,

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Dated: October 12, 2001

By:



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Appendix to the Amendment of October 12, 2001

**IN THE SPECIFICATION**

Please replace the first full paragraph on page 6 with the following paragraph:

This invention also provides a recombinant chromosome comprising a polynucleotide containing a nucleotide sequence, wherein the sequence includes at least one modification of the *NAP1L2*[*Napsll*] *Nap1l2* gene, wherein the modification is selected from a) substitution, b) deletion, c) frame-shift, d) aberrant insertion, e) altered epigenetic control, or f) site-directed mutagenesis that causes a loss of biological function in the *NAP1L2* gene.

**IN THE CLAIMS**

Please amend the following claims:

1. (AMENDED) A method for screening neural system defects in chromosomal material of a mammal, said method comprising:

[(A)] providing chromosomal material from said human;]

[(B)](A) detecting a modification of [the] a *NAP1L2* gene or a *Nap1l2* gene in the chromosomal material, wherein [said]the modification is selected from a) substitution, b) deletion, c) frame-shift, d) aberrant insertion [aberrant]or e) altered epigenetic control[:]  
that causes a loss of biological function in the *NAP1L2* gene or the *Nap1l2* gene; and

[(C)](B) correlating the modification of [said]the gene with a potential for a neural system defect.

2. (AMENDED) [A]The method [according to]of claim 1[where the said screening of neural system defects concerns a human being] wherein the mammal is a human.

3. (AMENDED) The method of claim 1, wherein [said] the modification in the NAP1L2 gene or the Nap1l2 gene is detected by hybridization with a labeled probe.

5. (AMENDED) [A]The method of claim 1, wherein [said] the modification is detected by

(A) amplification of the chromosomal material using PCR;

(B) sequencing [said] the chromosomal material to detect the modification of the nucleotide sequence; and

(C) correlating the modification of [said] the gene with a potential for neural system defects.

6. (AMENDED) A method for screening neural system defects in [a]human or mouse biological material, said method comprising:

[(A) providing biological material from said human;]

[(B)][A] detecting the absence, inappropriate, or modified expression of a NAP1L2 gene product or a Nap1l2 gene product using labeled antibodies to [said]the gene product; and

[(C)][B] correlating [said]the absence, inappropriate, or modified expression with a potential for neural system defects.

7. (AMENDED) The method of claim [5]6, wherein the [said]antibodies are polyclonal or monoclonal.

9. (AMENDED) [A]The method of [any one of claims]claim 1[to 8], wherein the neural system defect results from at least one of a failure of neural tube closure, [or]incomplete neural tube closure, inappropriate control of nucleosome activity in neurons, inappropriate control of the cell cycle in neurons, inappropriate differentiation of neurons, and inappropriate maintenance of neurons.

15. (AMENDED) A recombinant polynucleotide comprising a nucleotide sequence, wherein said sequence includes at least one modification of [the NAP1L2] a NAP1L2 gene or a Nap1l2 gene, wherein [said] the modification is selected from a) substitution, b) deletion, c) frameshift, d) insertion, or e) site-directed [mutagenis]mutagenesis that causes a loss of biological function in the NAP1L2 gene or the Nap1l2 gene.

17. (AMENDED) The polynucleotide of claim 15 [or 16], wherein said polynucleotide is a chromosome or a part of [it] a chromosome of a neural cell.

19. (AMENDED) The neural cell of claim 18, wherein the cell is derived from an immortal cell line, [such as embryonic stem cells], neuronal cell line, [or] tumor derived cell line, embryonic stem cell, or wild type animal.

20. (AMENDED) The neural cell of claim 18, wherein the NAP1L2 gene or the Nap1l2 gene is under control of a neural-specific promoter, such as nestin, other neuronal members, and inducible promoters.

22. (AMENDED) The neural cell of claim 18, [20, or 21,] wherein the NAP1L2 gene or the Nap1l2 gene is modified, wherein said modification is [selected from] a) substitution, b) deletion, c) frameshift, d) insertion, [or] e) site-directed mutagenesis or f)

naturally occurring mutation that causes a loss of biological function in the *NAP1L2* gene or *Nap1L2* gene.

24. (AMENDED) A polynucleotide comprising the promoter of a *Nap1L2* gene in SEQ ID NO:1, a polynucleotide hybridizing under stringent conditions with SEQ ID NO: 1, at least 20 nucleotides of SEQ ID NO: 1, the promoter of [the] a *NAP1L2* gene in SEQ ID NO: 4, [or] a polynucleotide hybridizing under stringent conditions with SEQ ID NO: 4, [or] at least 20 nucleotides of [said] SEQ ID NO: 4, SEQ ID NO: 6, a polynucleotide hybridizing under stringent conditions with SEQ ID NO:6, or at least 20 nucleotides of SEQ ID NO: 6.

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(SEQ ID NO. 6)

Genomic sequence BPX human

1. acctaaaggaaaaatttatctataaaactgacagaatttagaaataaatacaacaatatgttaacagcttttaatatctctg  
2. atagtcaacaattcttttaaatctggaataataatagtcacttaaaattttaaaaaattgttcaattataaatgatccaag  
3. ttagaaatgatgacaaaaataaacctccaccaataattactatagaggagaaatttaattactgcgaagcttccatcccta  
4. caaatacattatcaaatagtttaaccattctttaatgctgagatttagattatcccaattactcaaaagcatcaagc  
5. aaatgttatgatttctaagaataaaacataaacttccattttggcttttgratatatgtatatttctaacggctgttaaaag  
6. ccagcattaagaaggagaaagcagaagtcagatttgggactggggttatttataagccaggcaactgggttaattgtgggt  
7. aattgtctggtagttttactagtacgttagttgtataccatactagttttcatcacaggccctcattcgccccact  
8. gccatcggaacttctctctctccctccacaggaaatgttctcgagaatttttcaaccataaatacatatagcttctgtaaaaa  
9. taccgacaaacataatatagaattttaaataaactgacacgcccacctaagaccatcagtgctaattcctgggtgtttta  
10. atcttttgaagcgtttgtttatcagctcttccaccatccacctctccccctcccgagctcctaaatacaagagat  
11. gatttaggagtggtgggtgcttctgtctctctcattgttcgacatttagttacgtttctctgagctctcggaaagc  
12. ataaaagtataatctctgttaaaagtggatgaatgaactaatgaacgcaatgggattccagaaaaactctgccccgagt  
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28. aggaggaggaggaggaggaggagggagcttactggagaagagaaataaagaagaggagatccttaagggaatt  
29. cctgatttttggctaaactgttttaaaaaacggttgatacactcactcctttgatttaagaataatgatgagcctattctgaa  
30. gctcctgacagatattaaagtttaagctttcagatcctggcgagccctcagtttcacactagaatttcacttcaaaacca  
31. atgnaattttcaaaaatgaattgttgcacaaagacattatgtgctgaagtcaaaagctagcatattatgatccccatccctat  
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33. gaagaacacagaaacatcggaatcggggaacaaatccgaactgtaactgaagattttcccaaggaatttcattttcaatttt  
34. tctctcctcatggaatcacctcaaatggaaggatggaatgatgattttttacttggctacaaatttacctacttacata  
35. attccaagatcagattattttttctcagggtgatgactggaattctcagcaggaggggtagttagagaagtttaattgatgc  
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38. agaagccctgtgtctgtattttgtctttagttagtttagtttaaaacatattgtttcaaaaataaagaaaggttcaaaact  
39. aatttaattgaccttgagtttagtagtagaattgttttcaagaaatgtacactgtggttaaatgattttaaaccacttagat  
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45. tgttagtcagatttttcaaaagtgtgcacataattgttctgtgtaattgtgaagccataattacagtggttattctctc  
46. ttttcttattacatcattcattgaaagtgatcactttaccattttgaaaagataatttcgtgttctctcactgcaaaataa  
47. aaagaataaaaaatttcaga

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26. (AMENDED) A method of making a recombinant neural cell comprising:

[(A) providing a neural cell;]

[(B)] (A) modifying a NAP1L2 gene or a Nap1l2 gene or [the] a promoter of the NAP1L2 gene or a promoter of the Nap1l2 gene in [the] a neural cell, wherein said modification is selected from a) substitution, b) deletion, c) frame-shift, and d) insertion that causes a loss of biological function in the gene; and

[(C)] (B) selecting modified cells.

27. (AMENDED) A method of screening for therapeutic compounds comprising:

[(A) providing a cell of any one of claims 18 to 23;]

[(B)] (A) introducing a compound to be screened to the cell of claim 18 [a compound to be screened]; and

[(C)] (B) correlating a change in the proliferation of the cells with the activity of the compound.

28. (AMENDED) A method of screening for therapeutic compounds comprising:

[(A) providing a transgenic knockout animal containing the human NAP1L2 gene in its chromosomes.]

[(B)] (A) introducing a compound to be screened to [the] a transgenic knockout animal containing the human NAP1L2 gene in its chromosomes [a compound to be screened]; and



[[C]] (B) correlating a change in the development and maturation of the transgenic knockout animal nervous system with the activity of the compound.

33. (AMENDED) A vector containing the nucleic acid molecule of claim [32] 24.

34. (AMENDED) A recombinant neural cell comprising a vector [comprising] wherein the vector comprises the *Nap1/2* gene or the *NAP1L2* gene.

35. (AMENDED) The neural cell of claim 34, wherein the *Nap1/2* gene or the *NAP1L2* gene is under the control of a neural-specific promoter, such as nestin or other neuronal genes or inducible promoters.

36. (AMENDED) A recombinant neural cell of claim 35, wherein the *Nap1/2* gene or the *NAP1L2* gene of the native cell is modified, wherein said modification is selected from a) substitution, b) deletion, c) frame-shift, d) insertion, or e) site-directed mutagenesis that causes a loss of biological function in the *Nap1/2* gene or the *NAP1L2* gene.

37. (AMENDED) A method of screening for therapeutic compounds in a cell comprising the nucleotide of claim 24, wherein the method comprises:

[(A) providing a cell containing a polynucleotide according to claim 32;]

[(B)] (A) introducing to the cell a compound to be screened; and

[(C)] (B) correlating a change in the proliferation of the cell with activity of the compound.

39. (AMENDED) [Use of therapeutic compounds obtained by the method according to claim 27 for] A method of increasing the expression of *NAP1L2* gene in

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tumoral human neural cells or for decreasing the expression of *NAP1L2* gene in human neural cells afflicted by a degenerating disease, comprising administering the therapeutic compounds of claim 27 to achieve an increase in expression of *NAP1L2* gene in tumoral human neural cells or a decrease in expression of *NAP1L2* gene in human neural cells afflicted by a degenerating disease.

41. (AMENDED) [A] The plasmid [consisting in the deposit made at] deposited at C.N.C.M. under the Accession Number I-2463, I-2464, I-2465, or I-2466.

45. (AMENDED) A polynucleotide [containing] comprising the sequence SEQ ID NO: 6.

46. (AMENDED) The polynucleotide of claim 24, wherein said polynucleotide further comprises [an] a heterologous amino acid sequence coding for [an] a heterologous polypeptide under the control of the *NAP1L2* promoter or the *Nap1l2* promoter.

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